

Figure 1

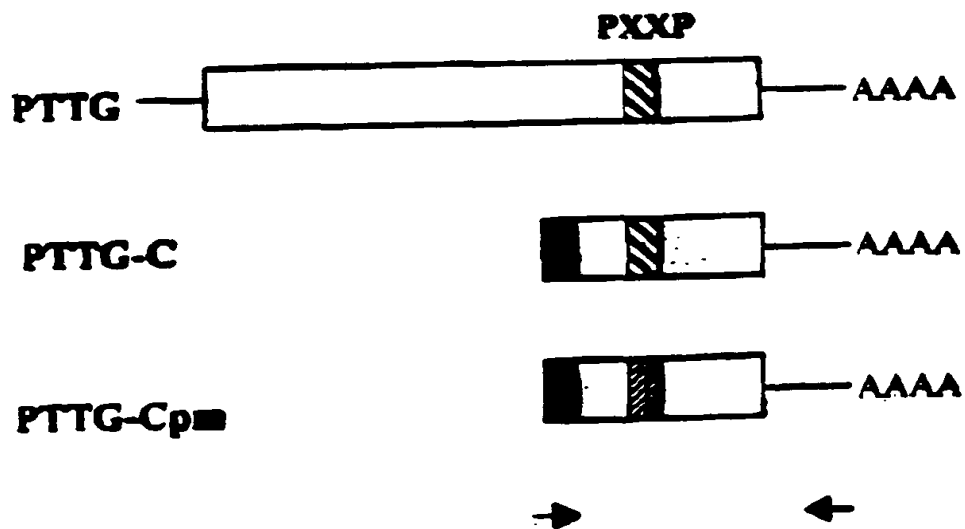


Figure 2A

09854326 051101

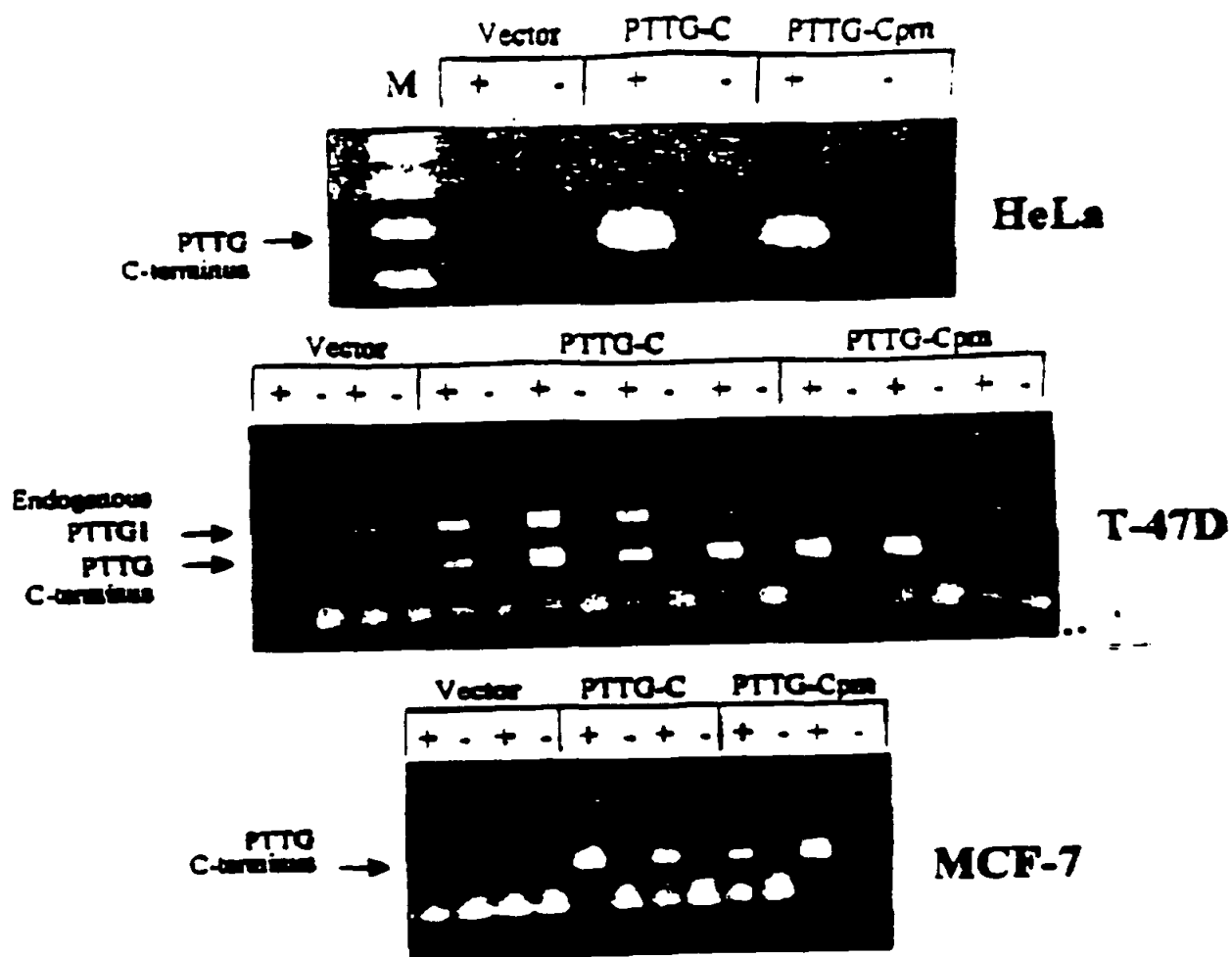


Figure 2B

09854326 051101

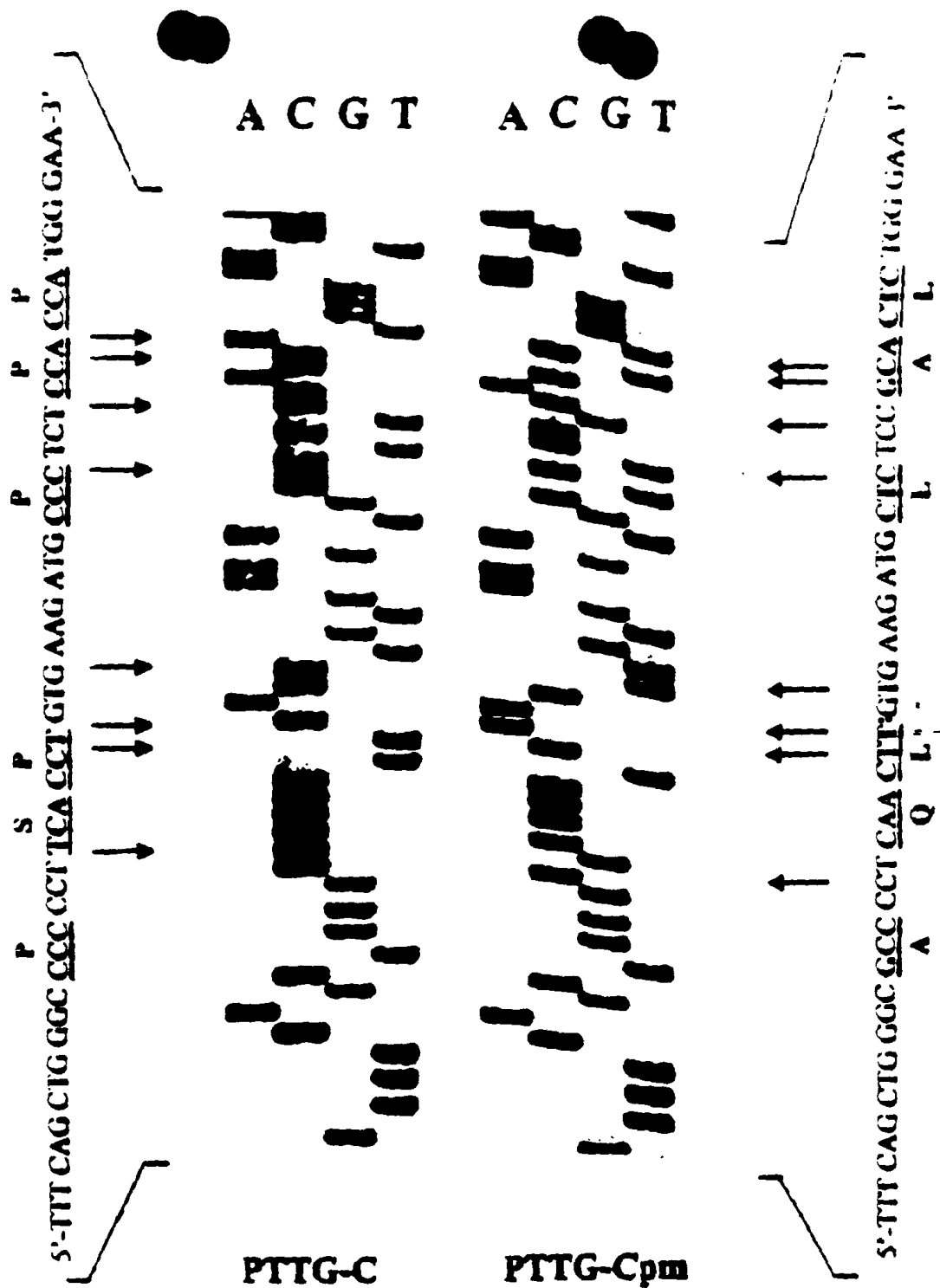


Figure 2C

0054326 .051101
T07T50" 92E45B60

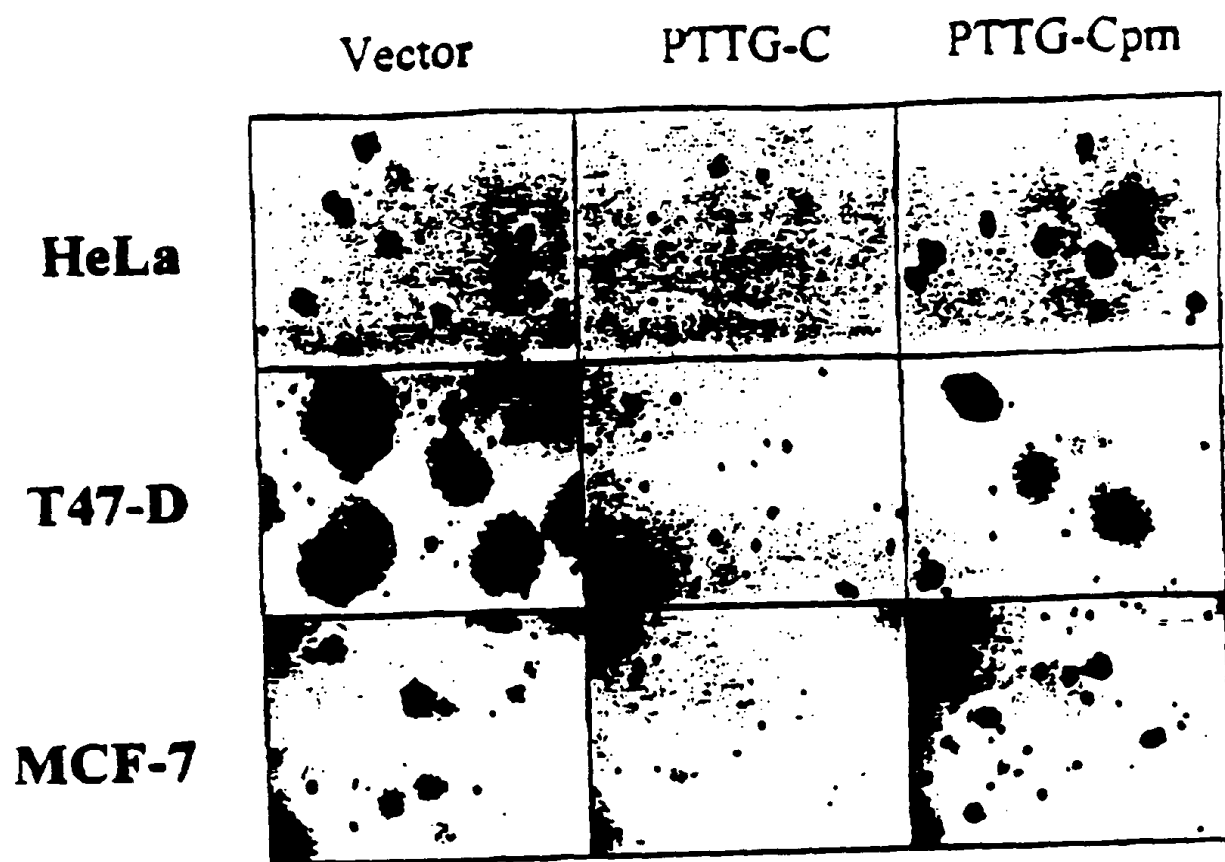


Figure 3

00554325 05101
T01F50" 92E45860

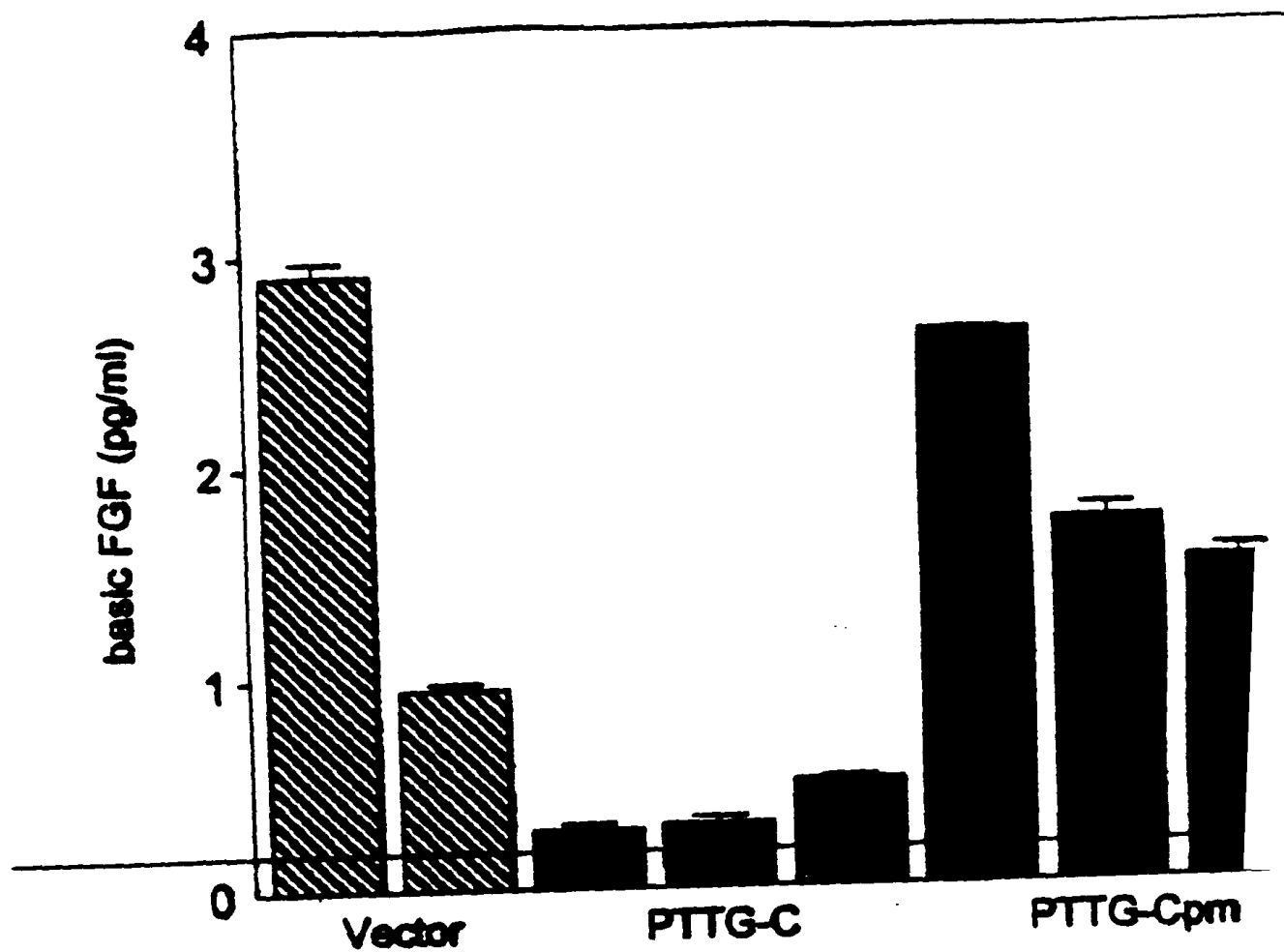
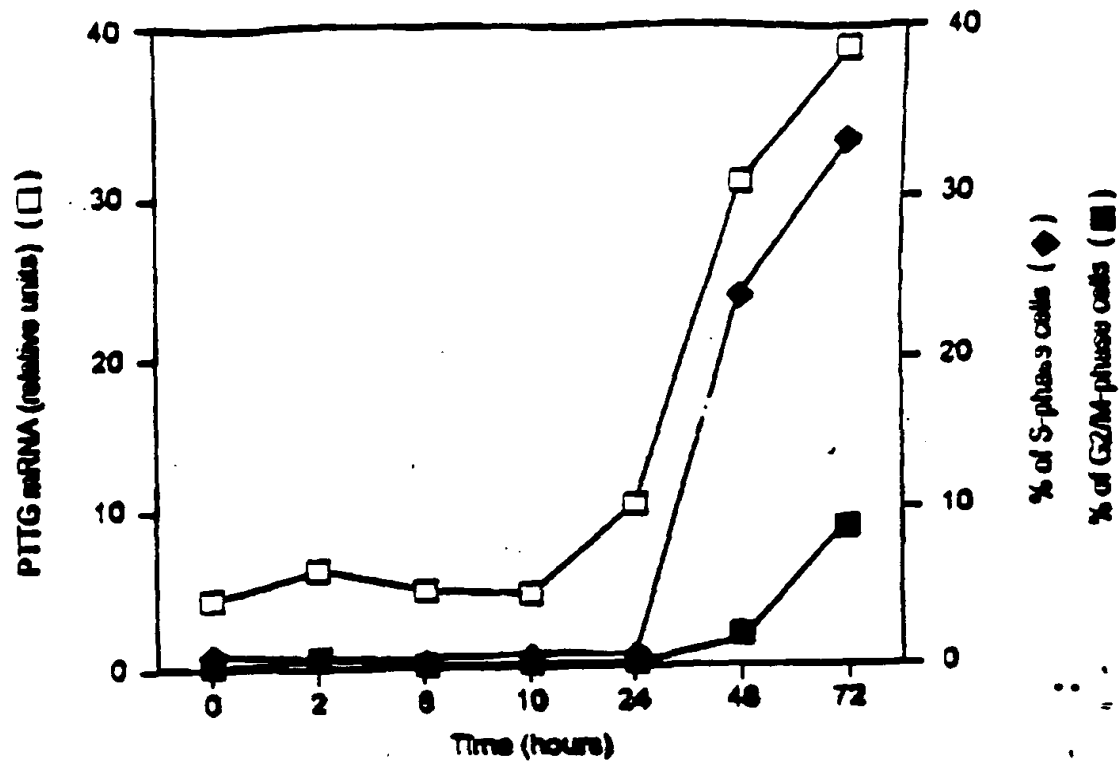


Figure 4.



pttg →



28S →

18S →

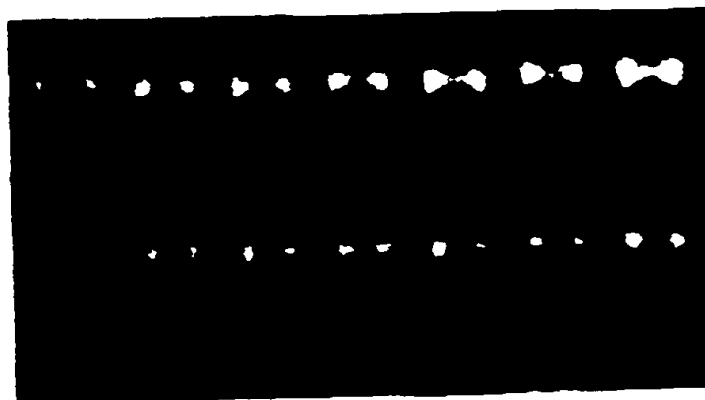


Figure 5

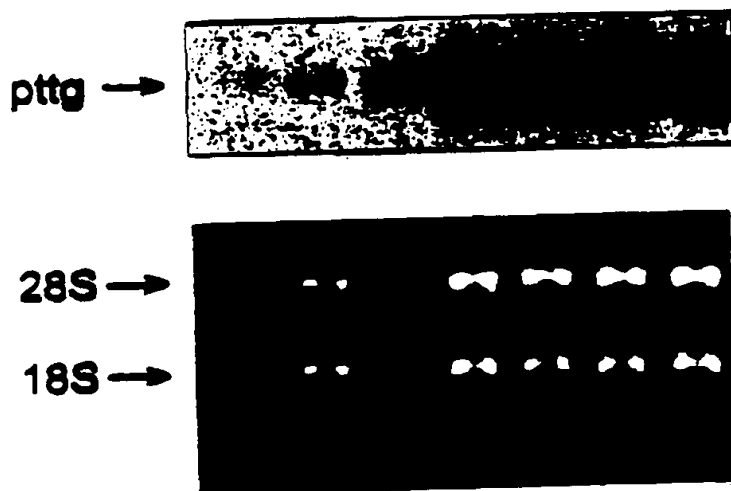
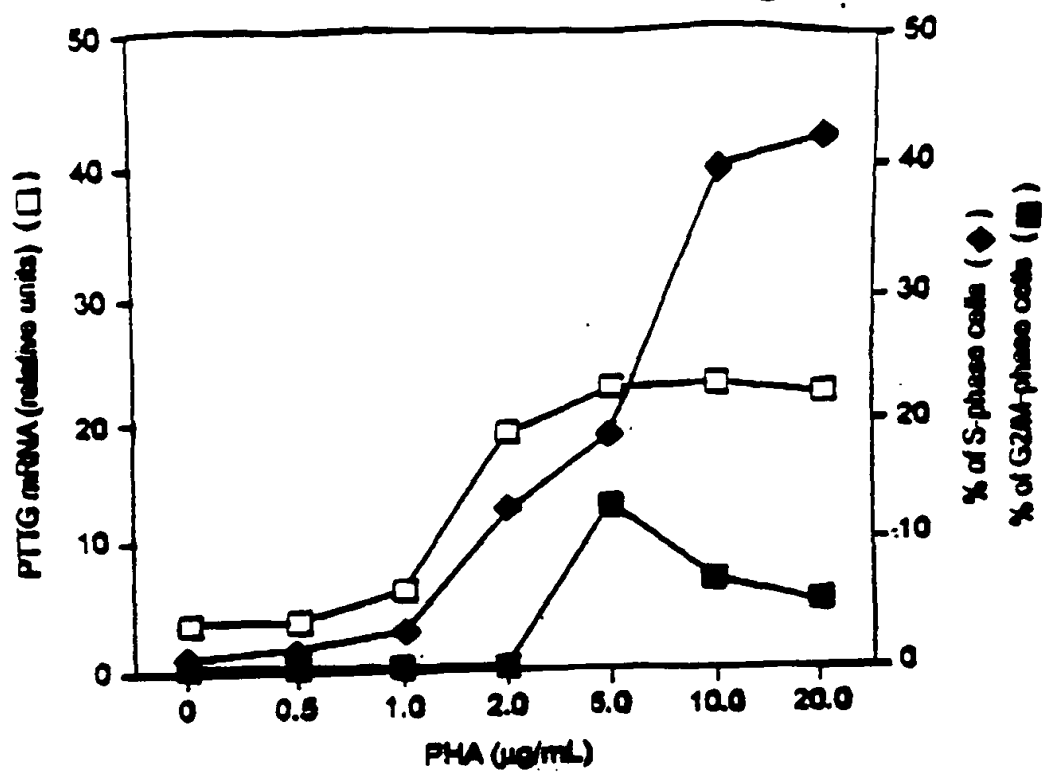


Figure 6

0954326-05101

Q3

IL-2

28S →

18S →

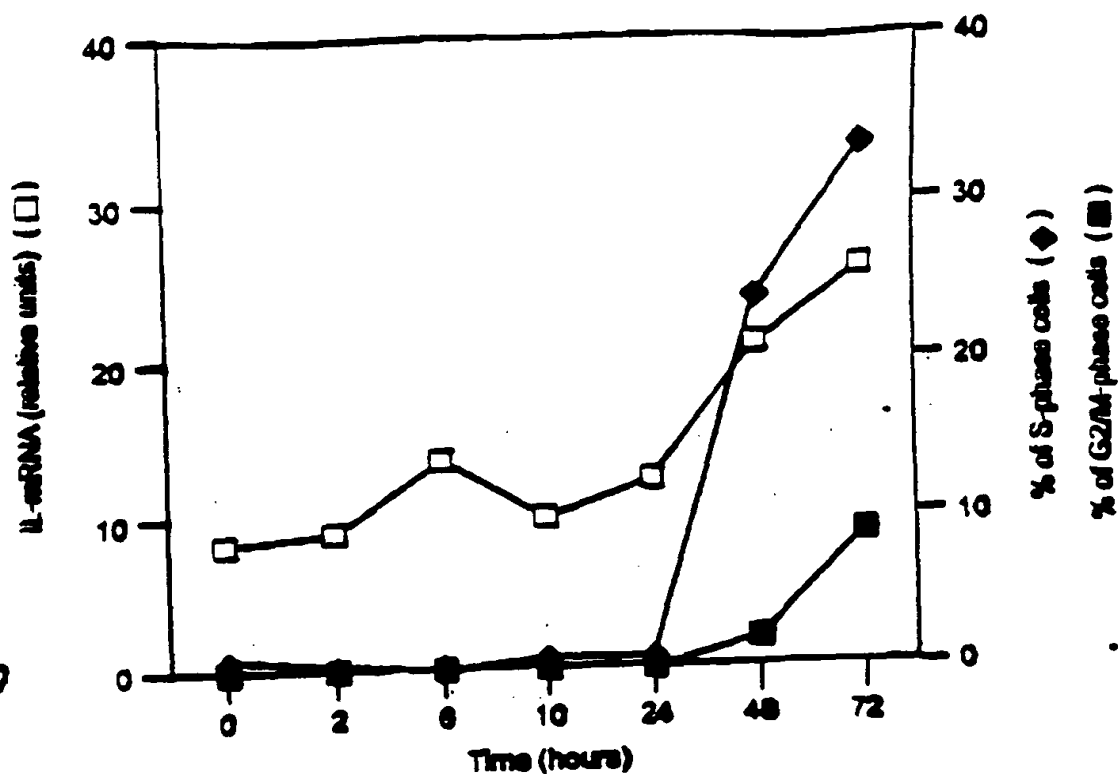


Figure 7

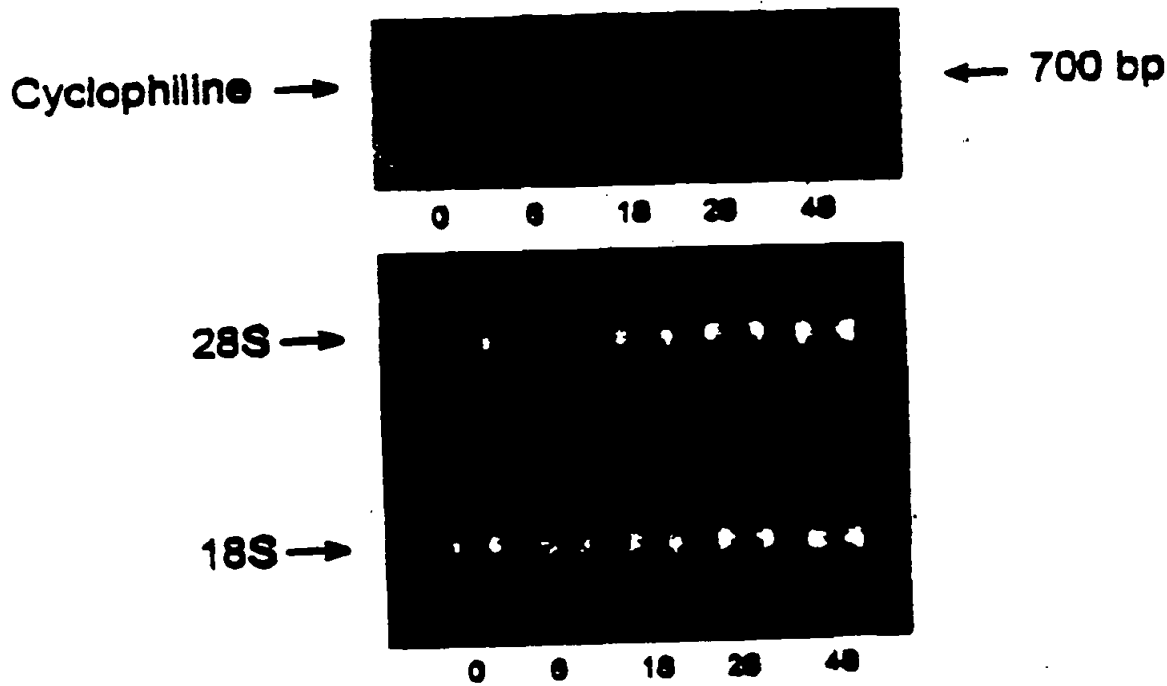
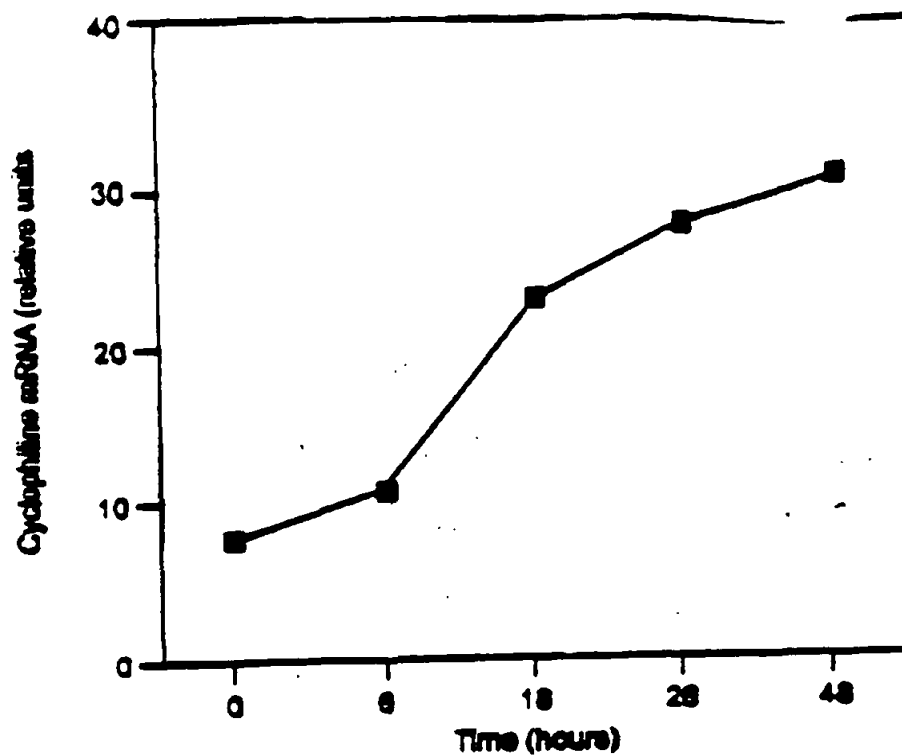


Figure 8

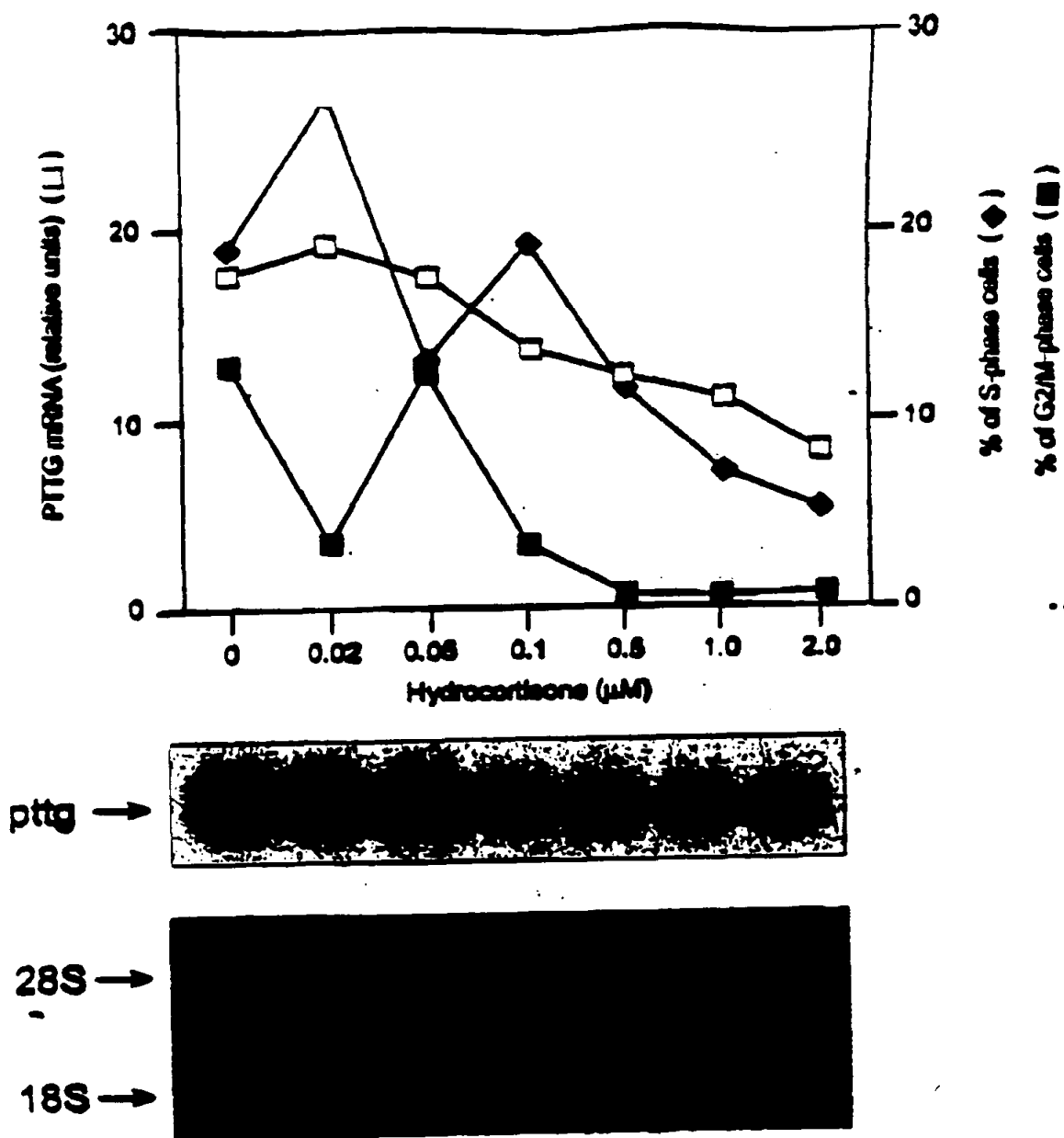


Figure 9

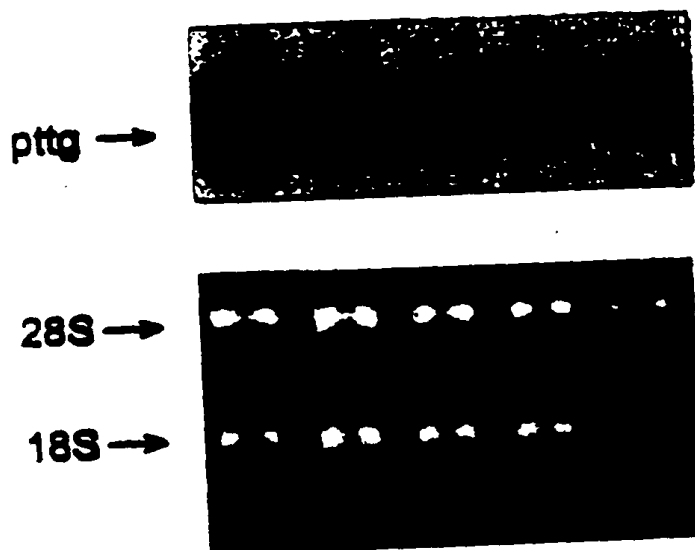
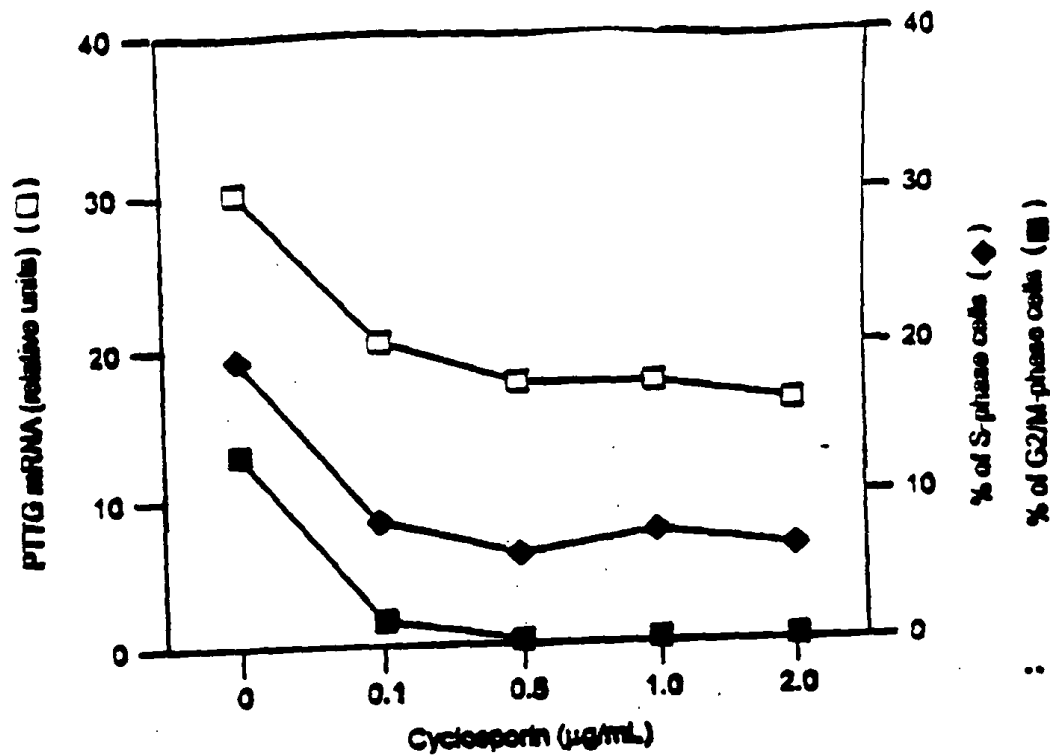
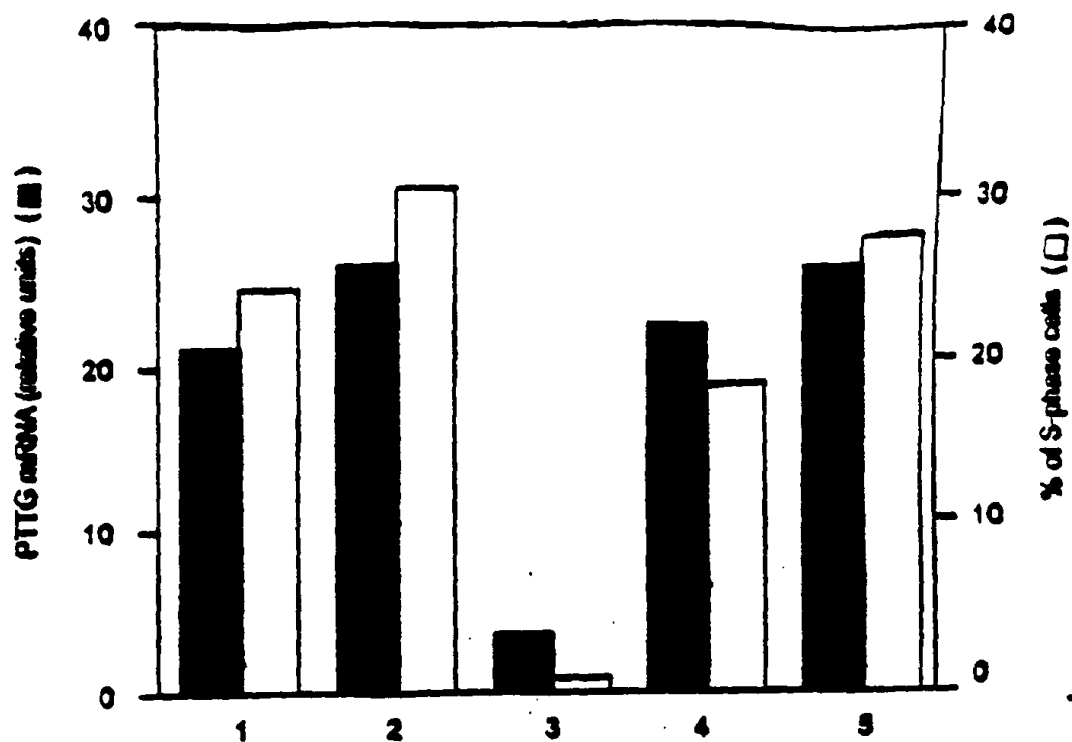


Figure 10

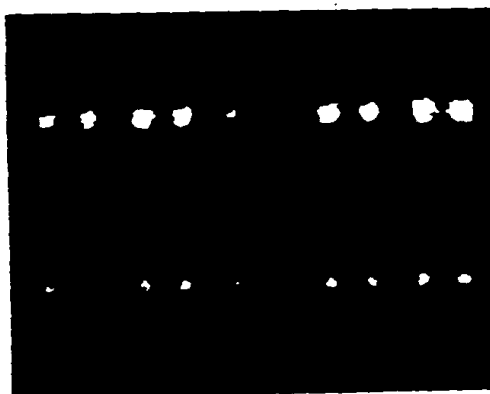
09854326.051101



pttg →



28S →



18S →

Figure 11

0545360 000000

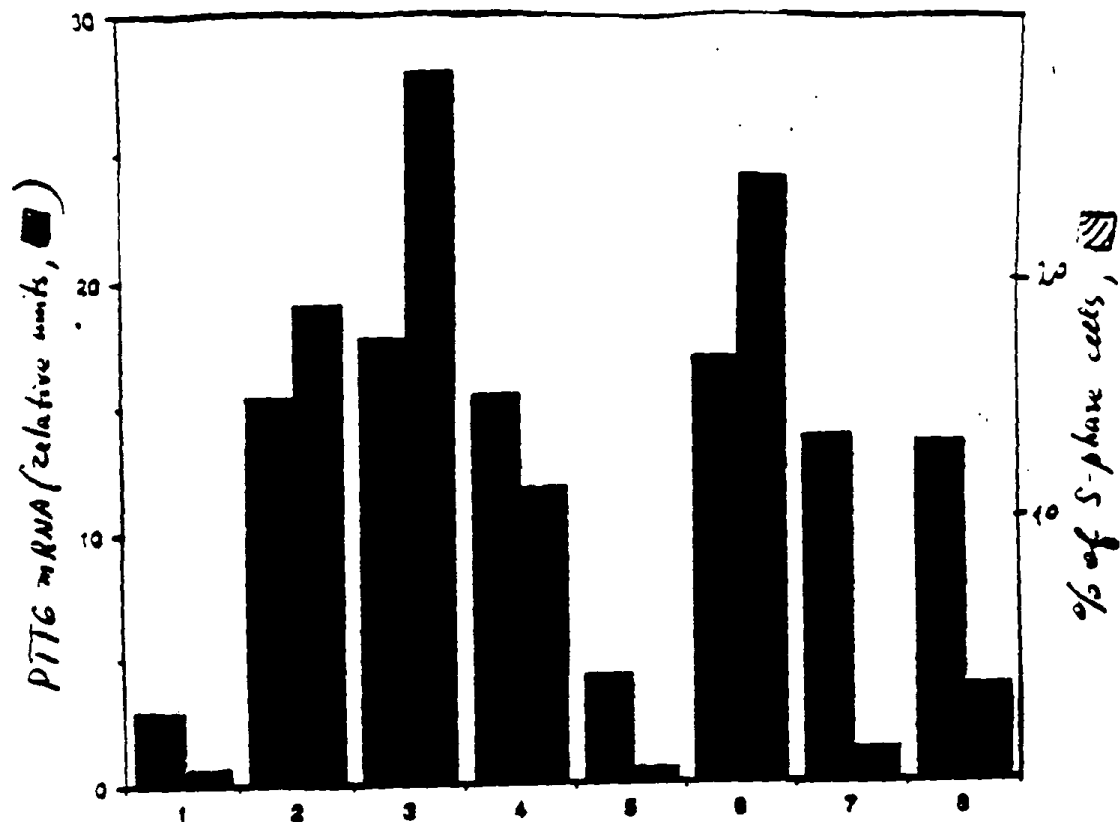


Figure 12

09854326 051101

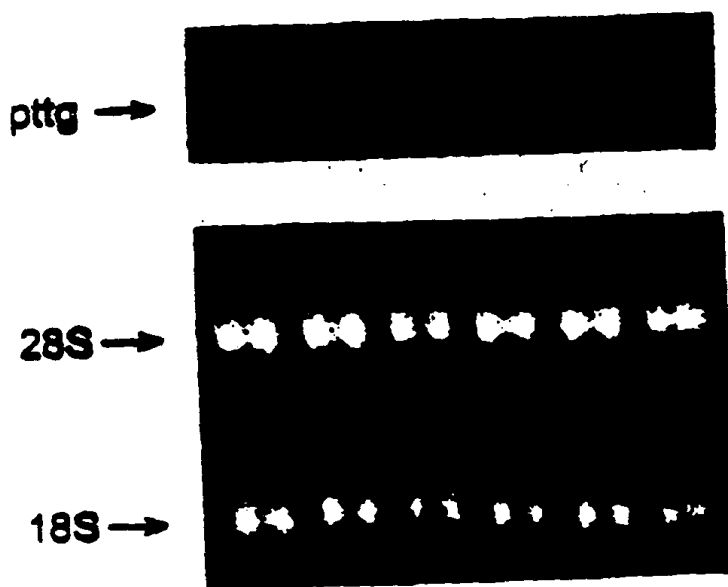
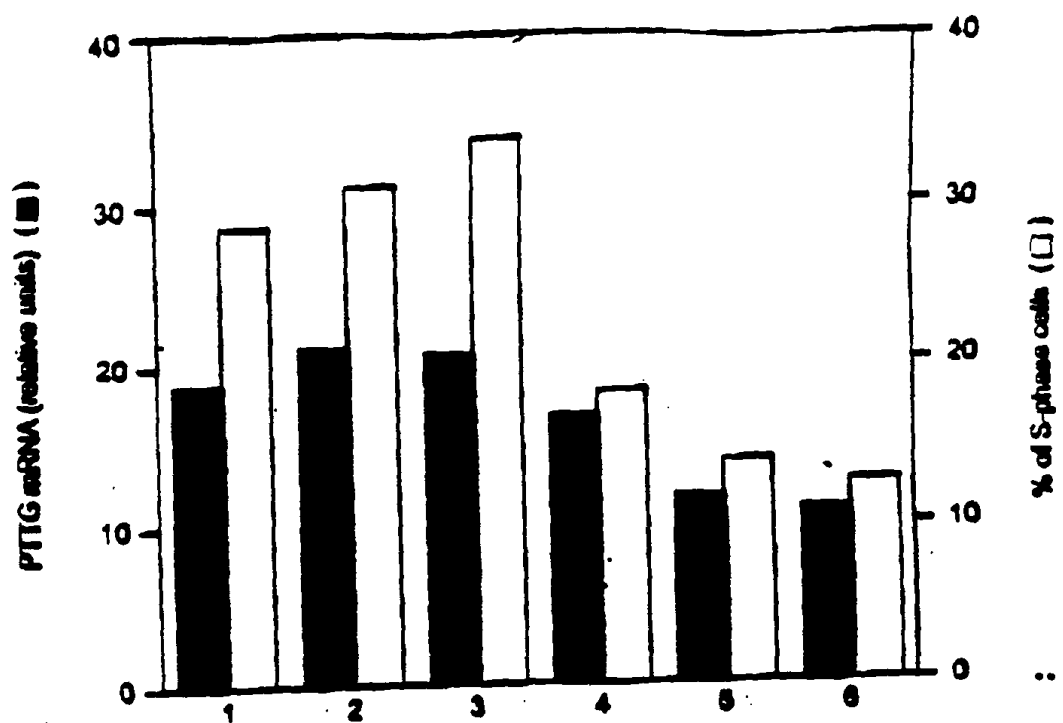


Figure 13

Figure 15

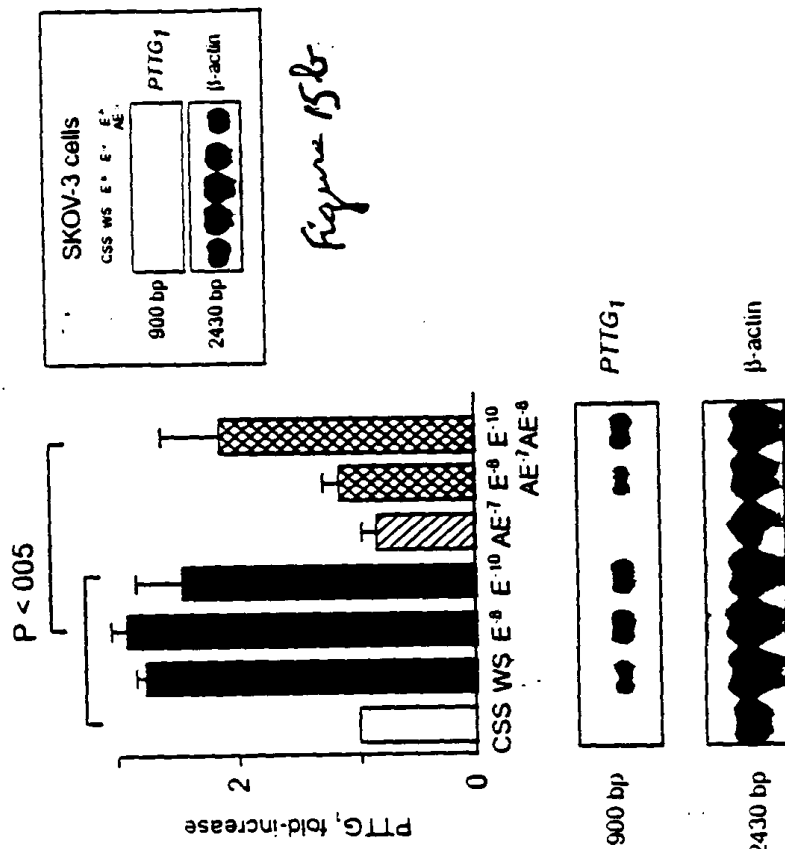


Figure 15b

Figure 15a

092432.05101
TOTSO" 92E45850

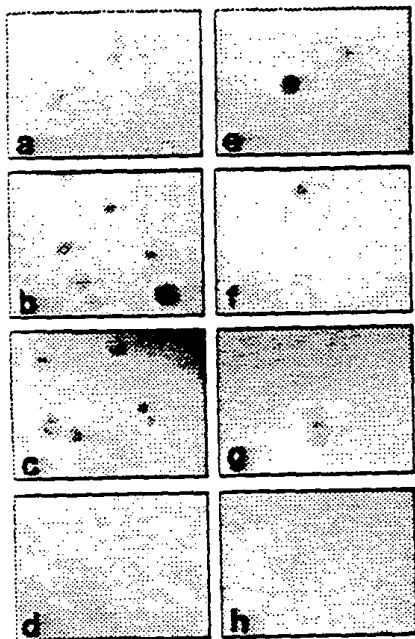


Figure 16

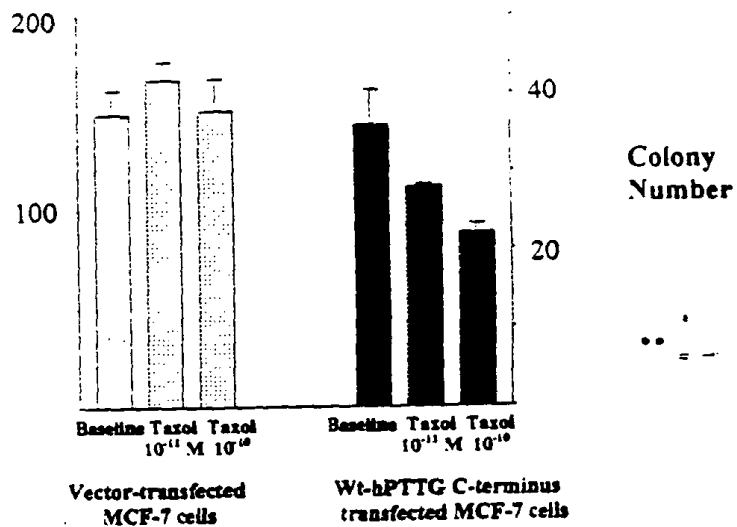


Figure 17

09245860

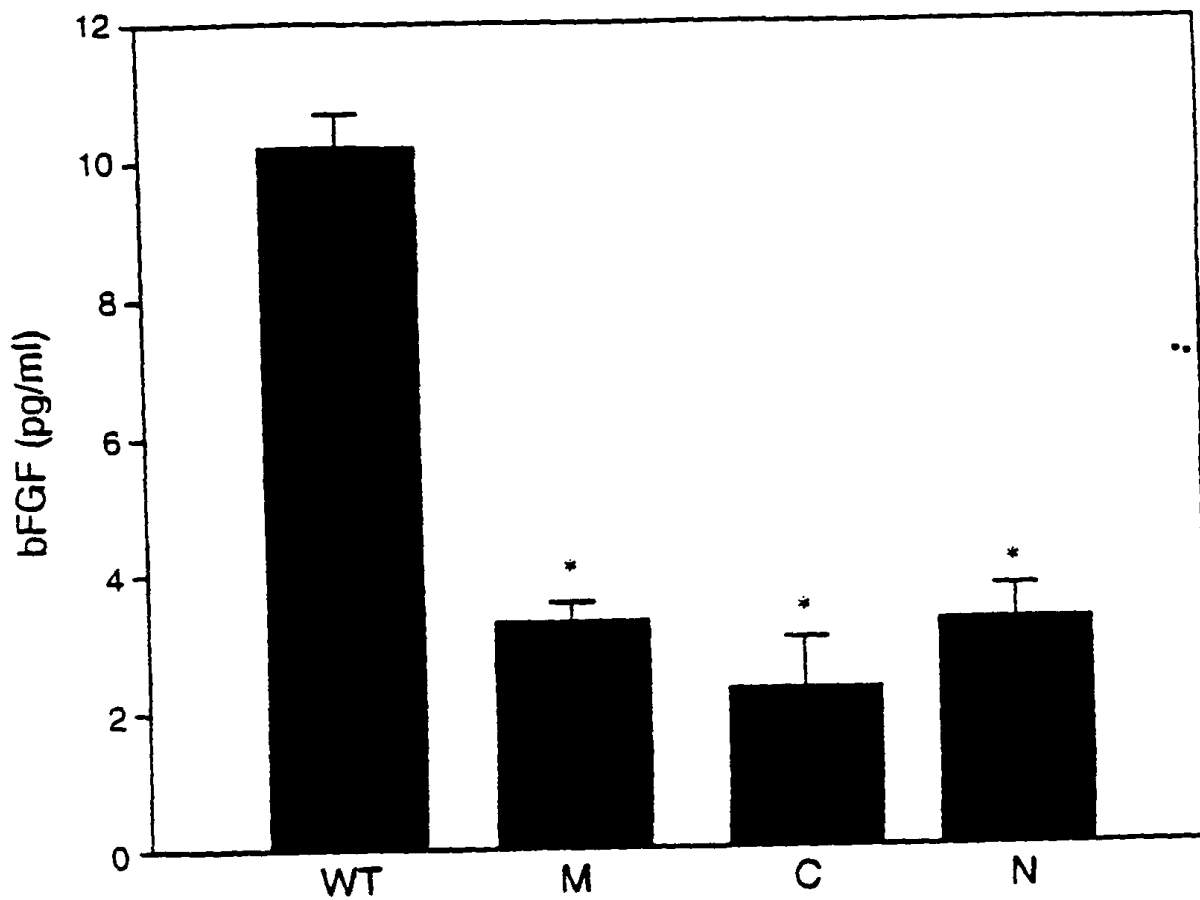


Figure 18

0954376 051101 10150 92545360

Fig. 20A

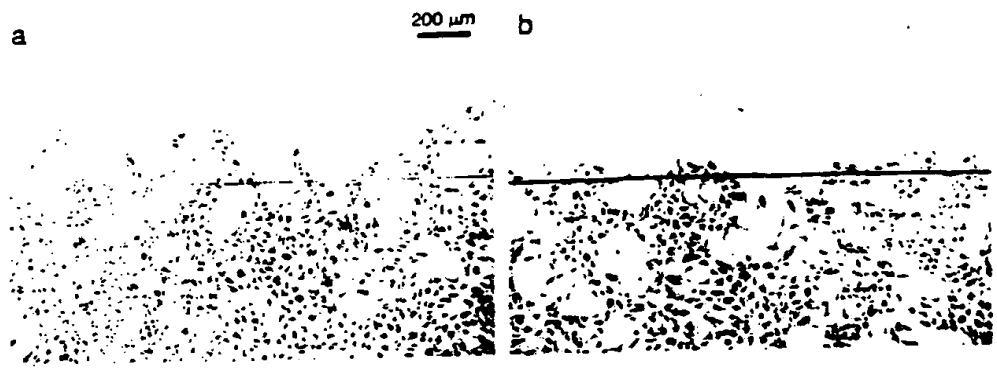


Fig. 20B

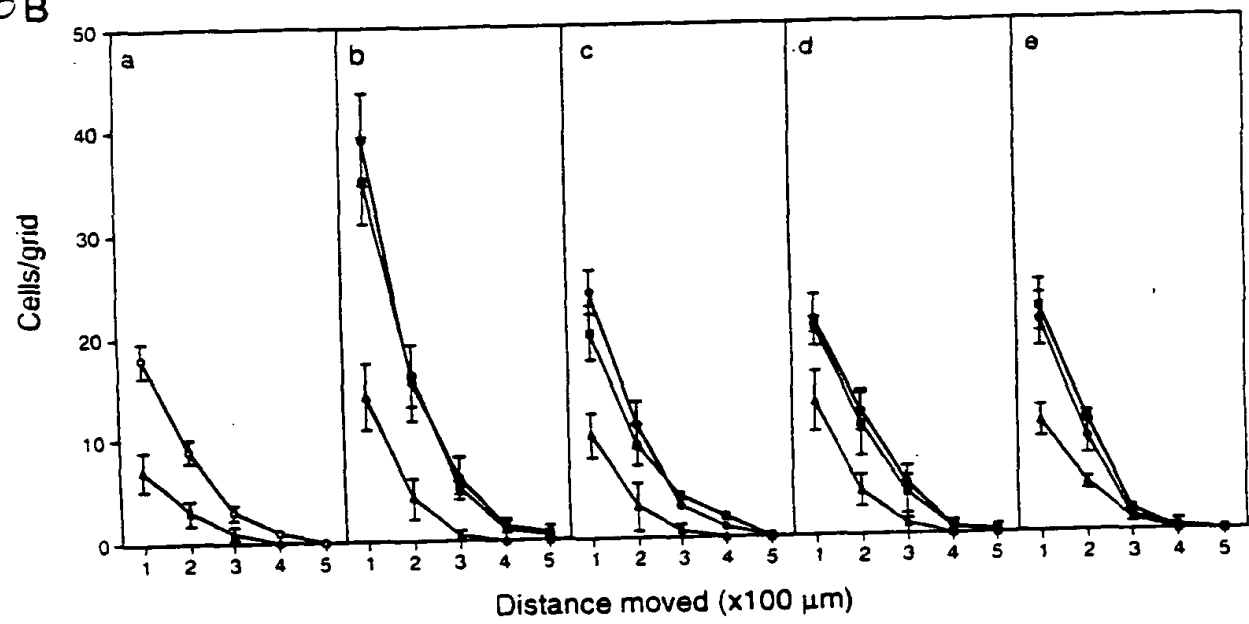


Figure 20

Fig. 21A

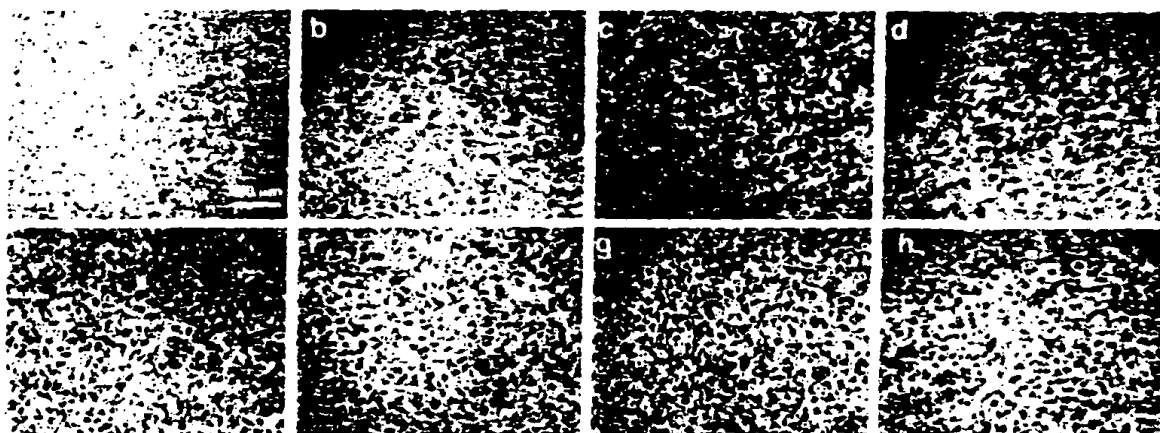


Fig. 21B

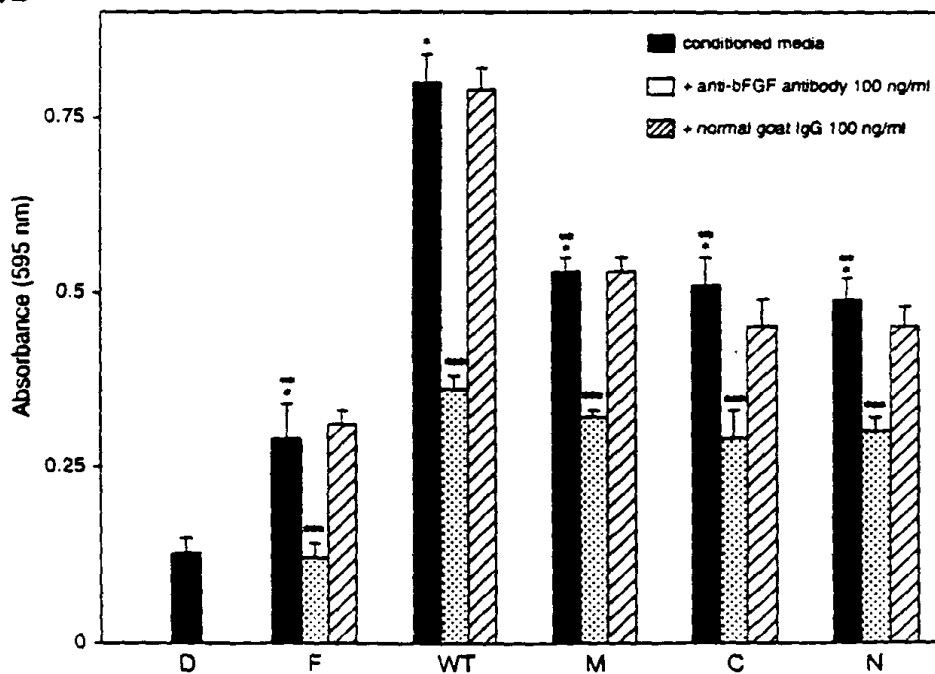


Figure 21

00854326 05101

0054325 051101

Fig 22A

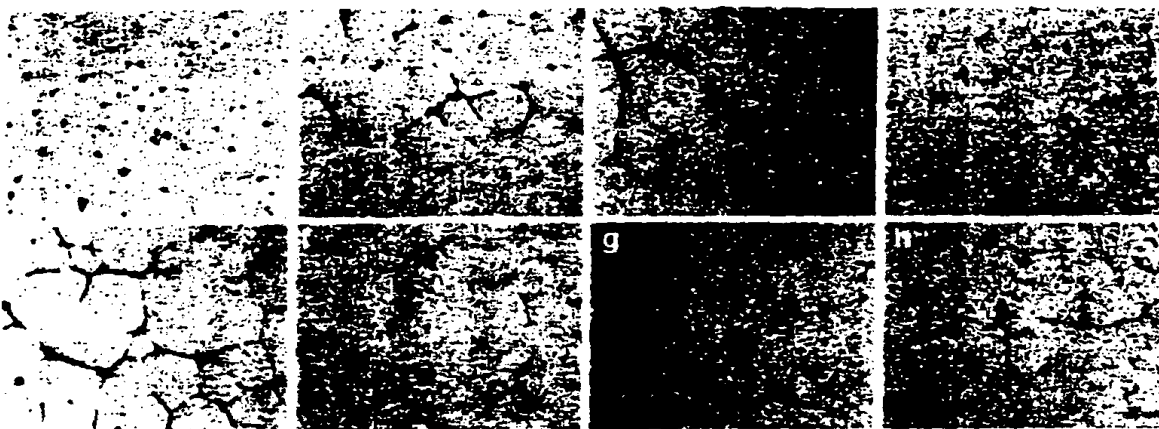


Fig 22B

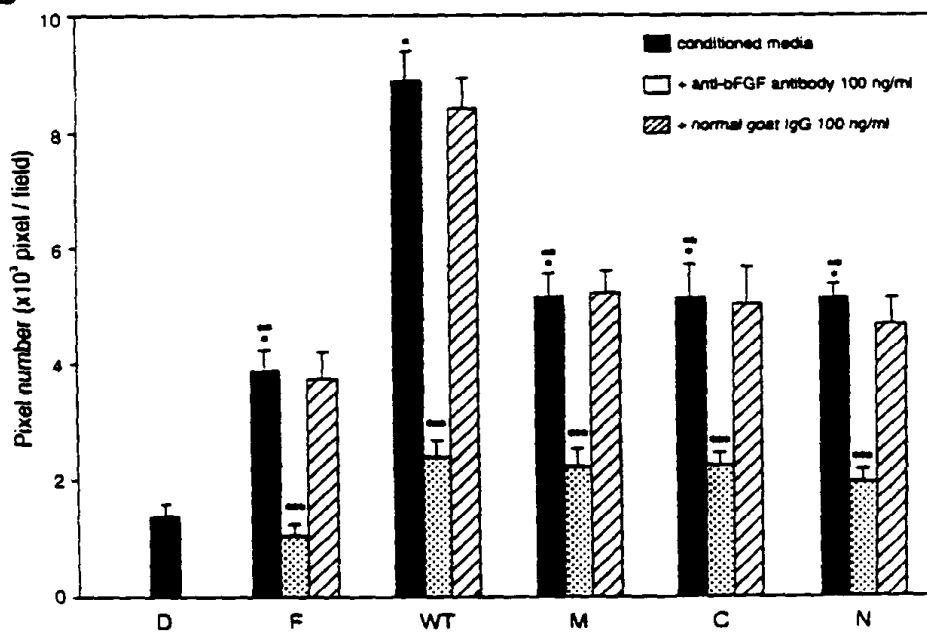


Figure 22

Fig 23A

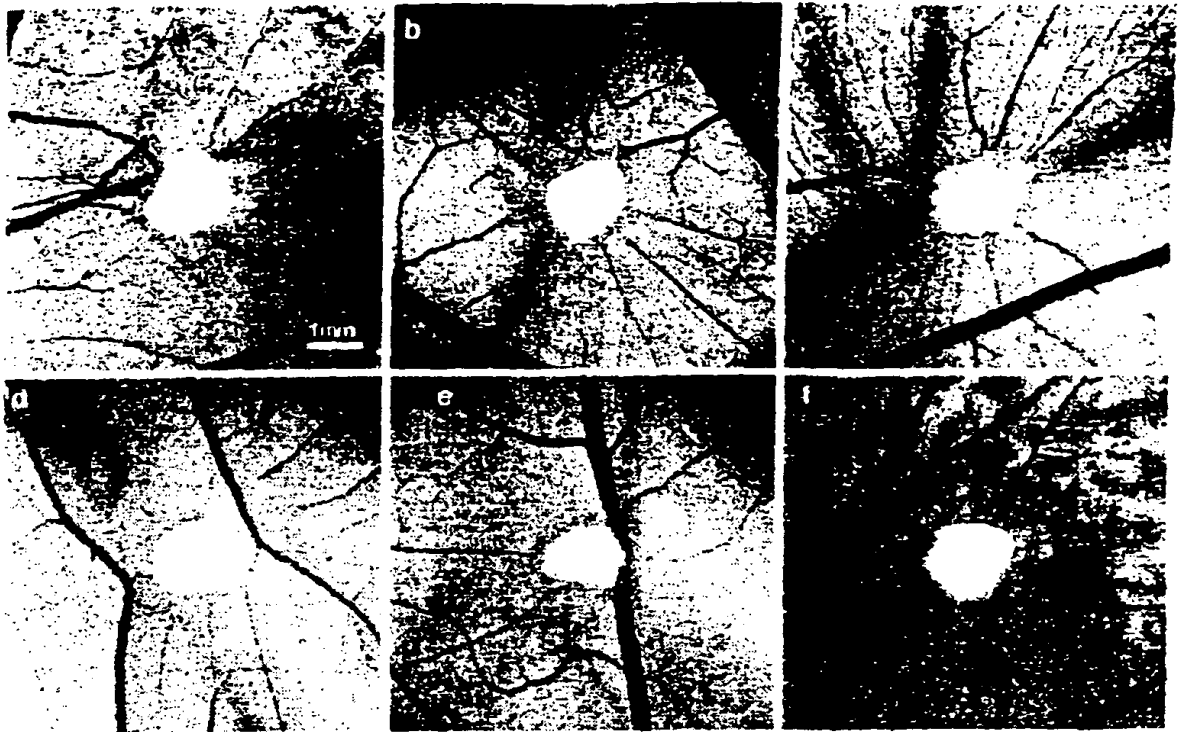


Fig. 23B

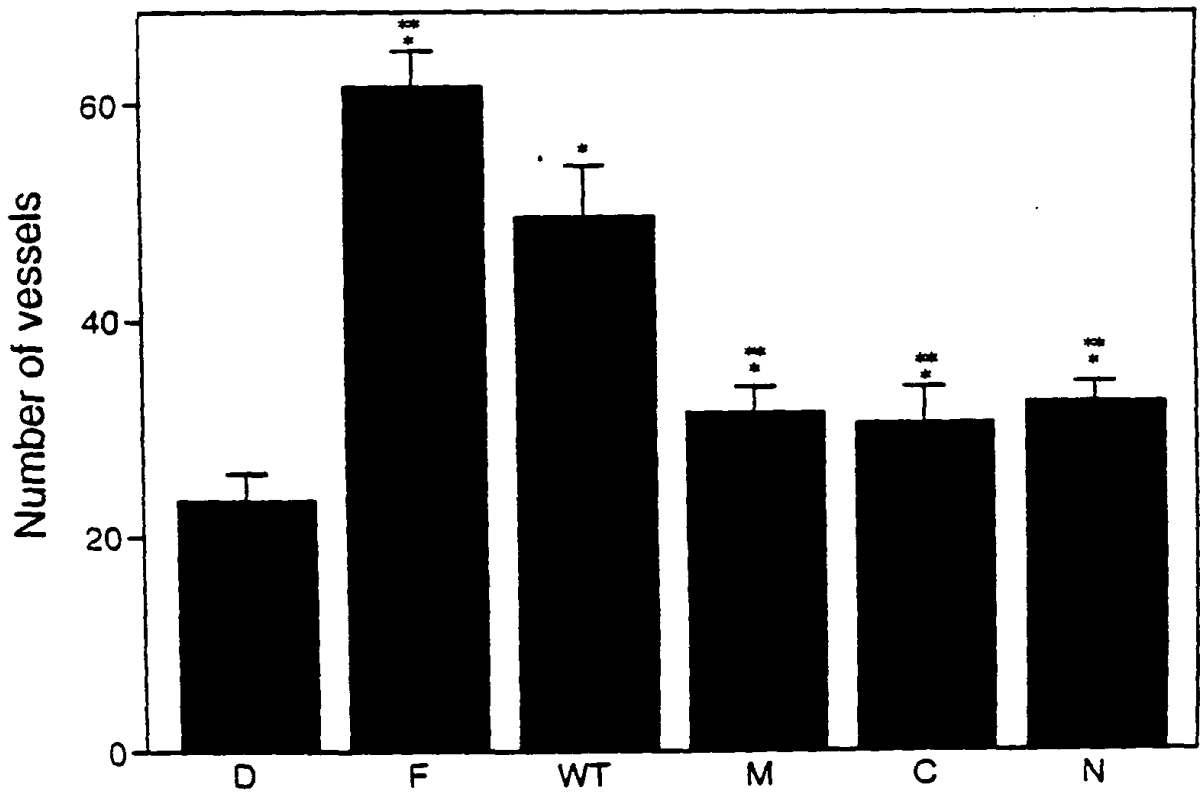


Figure 23

hPTTG1	1	MATLIYVDKE	NGEPGTRVVA	KDGLKLGSGP	SIKALDGRSQ	VSTPRFGKTF	DAPPALPKAT
hPTTG2	1	*****	I*****	**V**E**R*	*****I**	*L*****Y	***S*****
hPTTG3	1	*****	*E***IL*AT	*****E**R*	*****I**	***ISC*****	***TS*****
mPTTG	1	*****F****D	*E***R*LAS	*****T*V	*****KL*	*****V**V*	N*-**V***S
rPTTG	1	*****F****D	*E***S*LAS	*****V**V	*****KL*	*****V**V*	G*-*GL***S
	61	<u>RKALGTVNRA</u>	TEKSVKTKGP	LKQKQPSFSA	-KKMTEKTVKA	KSSVLASDDA	YPEIEKFFPF
	61	*****	*****	R*****	*****T	***P*****	*****L***
	61	*****	*****	*****	*****	*N**P*****G	*****L***
	58	*****V	A**PM**GK*	*QP***TLTG	**I***ST*T	Q***P*P***	*****L***
	58	*****V	**P**SSK*	*QS***TL*V	**I***ST*T	QG*AP*P***	*****L***
	121	NPLDFESFDL	PEEHQIAHLP	LSGVPLMILD	EERELEKLFQ	LGPPSPVKMP	SPPWESNLLQ
	121	*L*****	***R*****	*****	**G*****	***P*****	***C**FA
	121	*G*****	*****	*****	*****	*****L***	*****K*****
	118	***--***	*****SL**	*N*****IT*N	**G*****LH	*****L*T*	FLS***DP*Y
	118	D*****	*****SL**	*N*****N	**G*****LH	*D*****LQK*	FL***DP*P
	181	SPSSILSTLD	VELPPVCCDI	DI			
	181	VSFKHSVDPG	C				
	181	**L***L***	*****S**	**			
	175	**P*A*****	*****Y*A	**			
	178	**P*A**A**	*****Y*A	**			

(Seq. ID No. 4)
 (Seq. ID No. 64)
 (Seq. ID No. 67)
 (Seq. ID No. 14)
 (Seq. ID No. 2)

Figure 24

:

PTTG1 cDNA

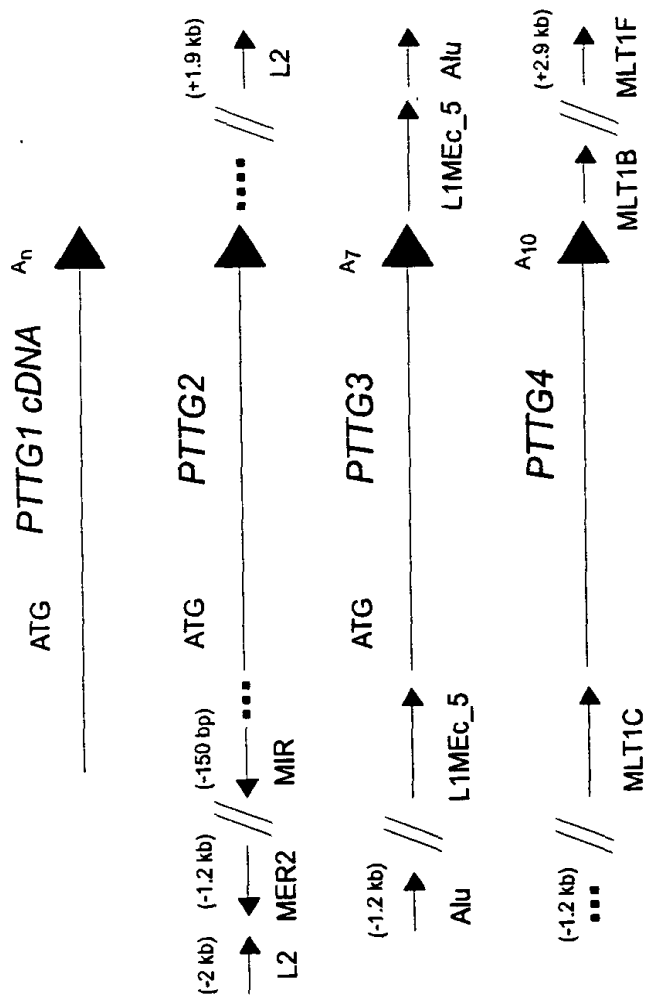


Figure 25

09854326 05110 92345860

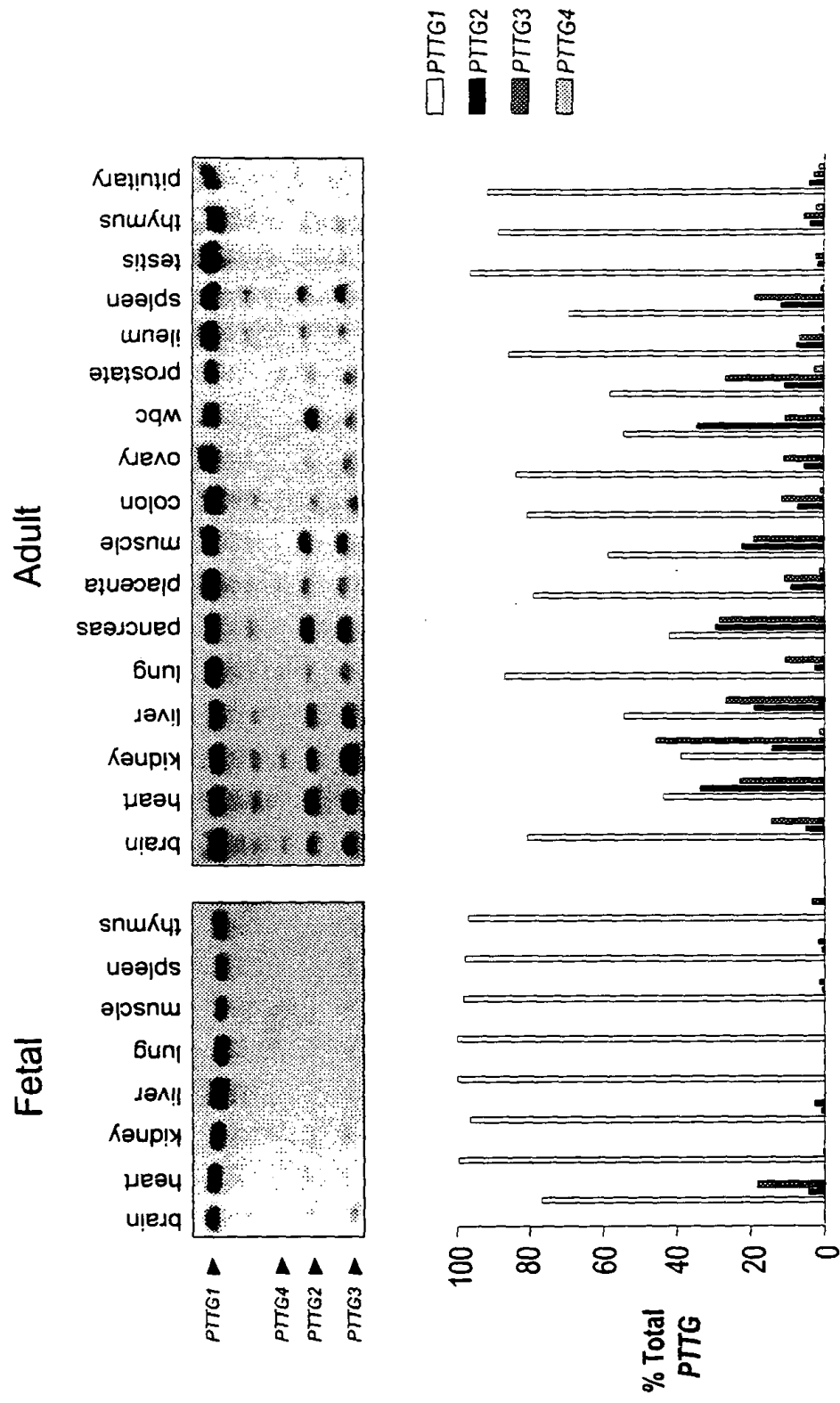


Figure 26 :

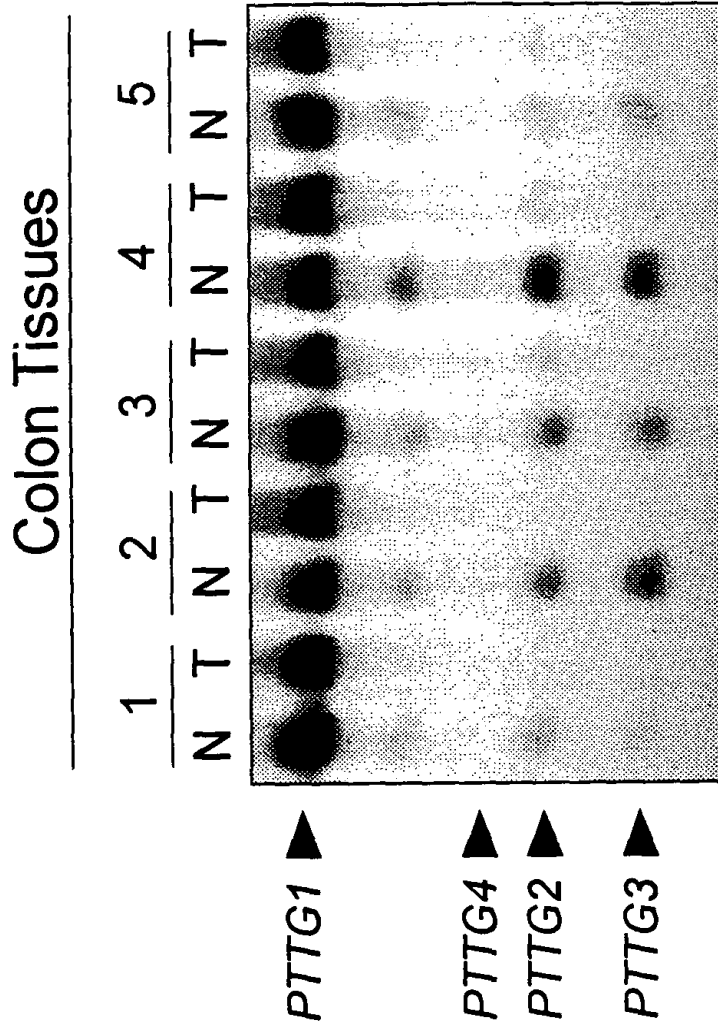


Figure 27A

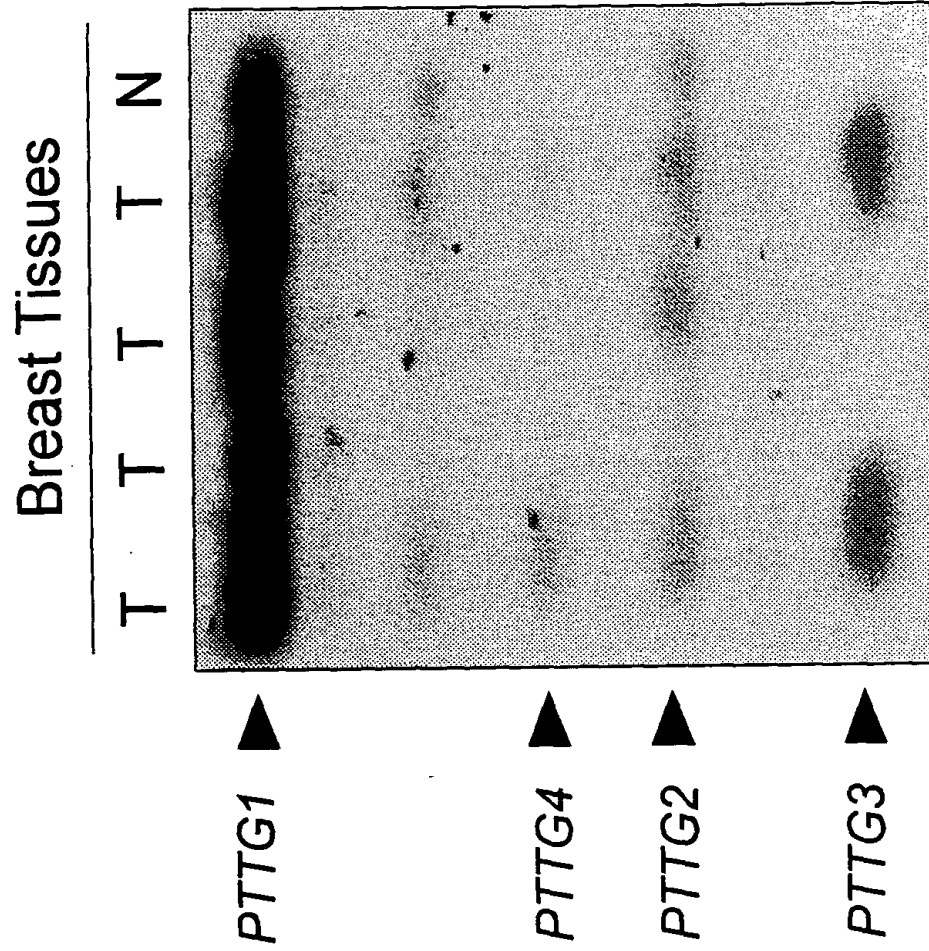


Figure 27B

Ovarian Tissues

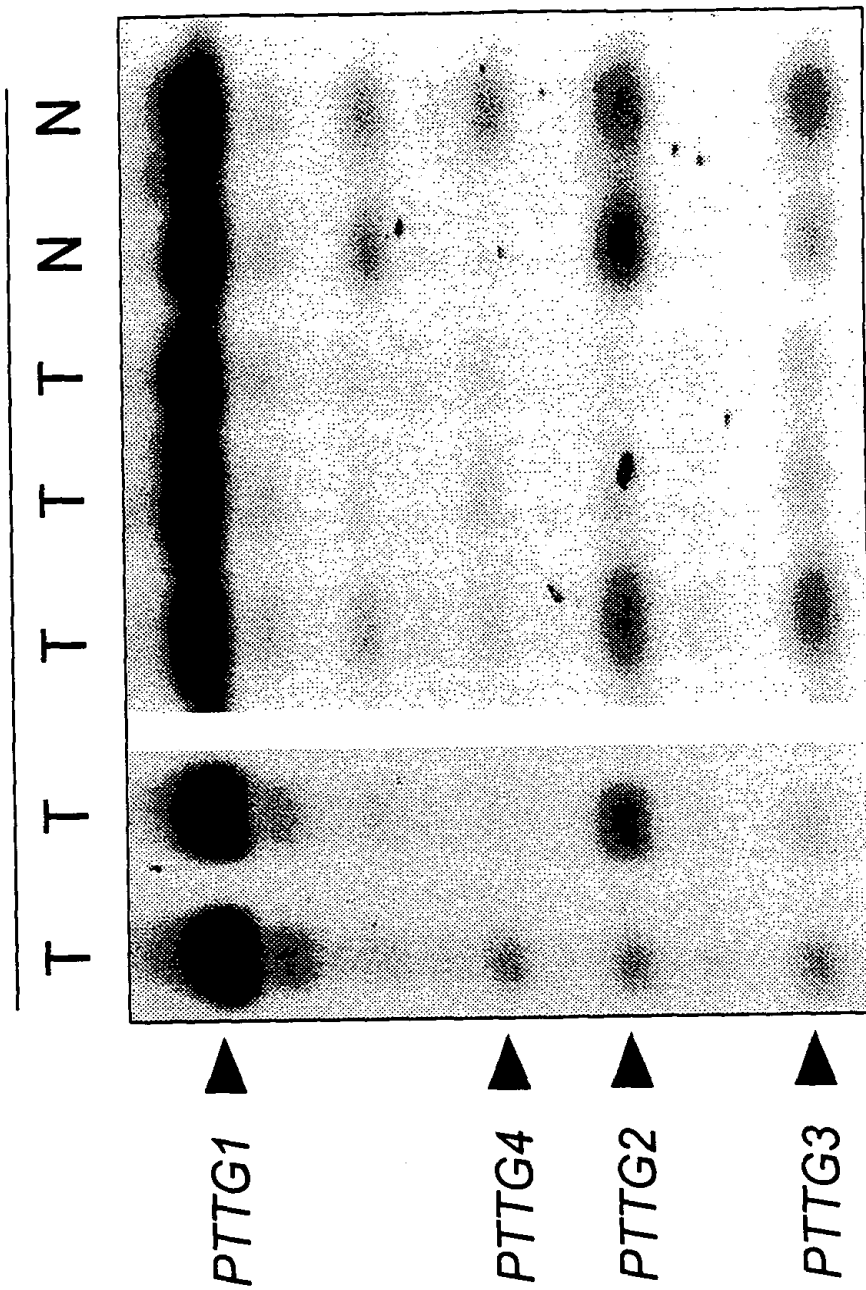


Figure 27C

09854326 05.11.01

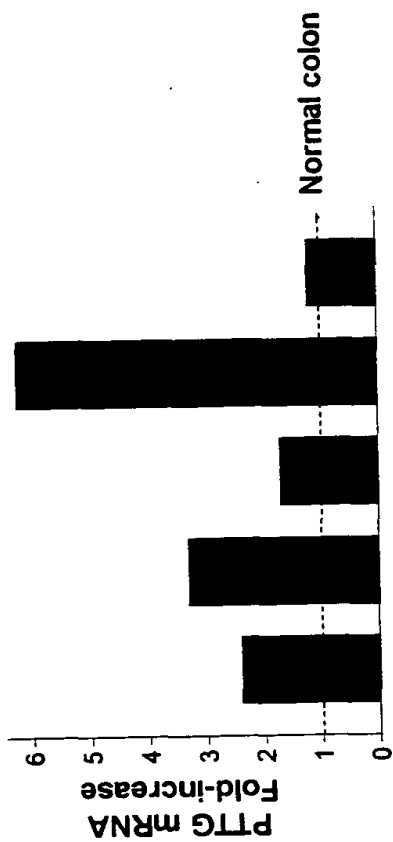


Figure 28

TOTTSO" 92E45860

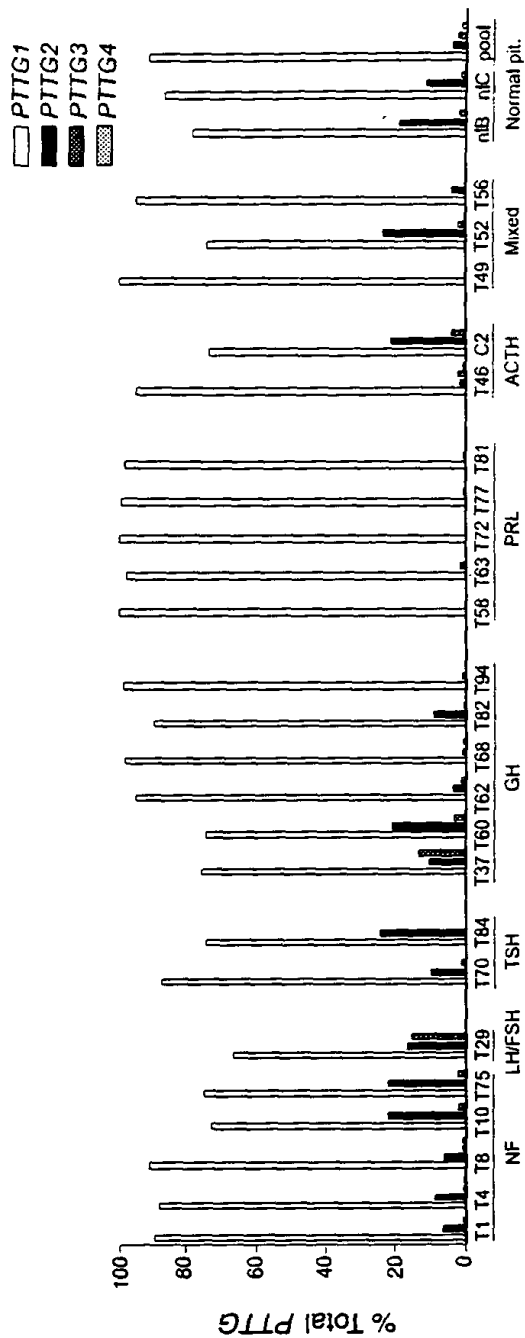


Figure 29

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Interspecific hybrids

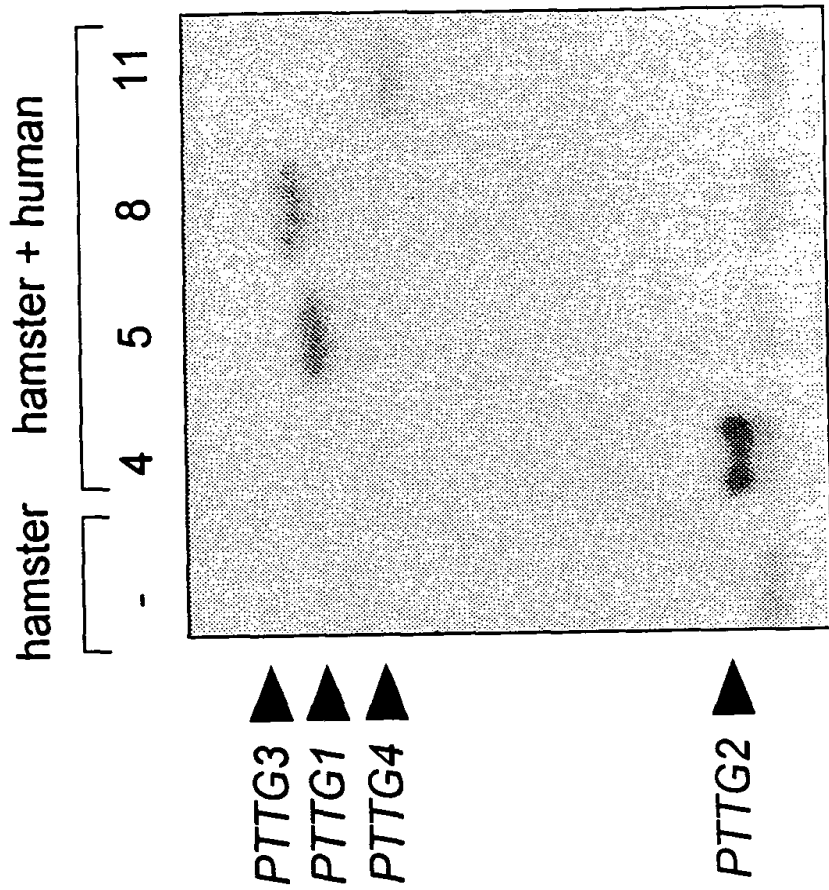


Figure 30A

TTT50" 92EH5860

Colon Tissues

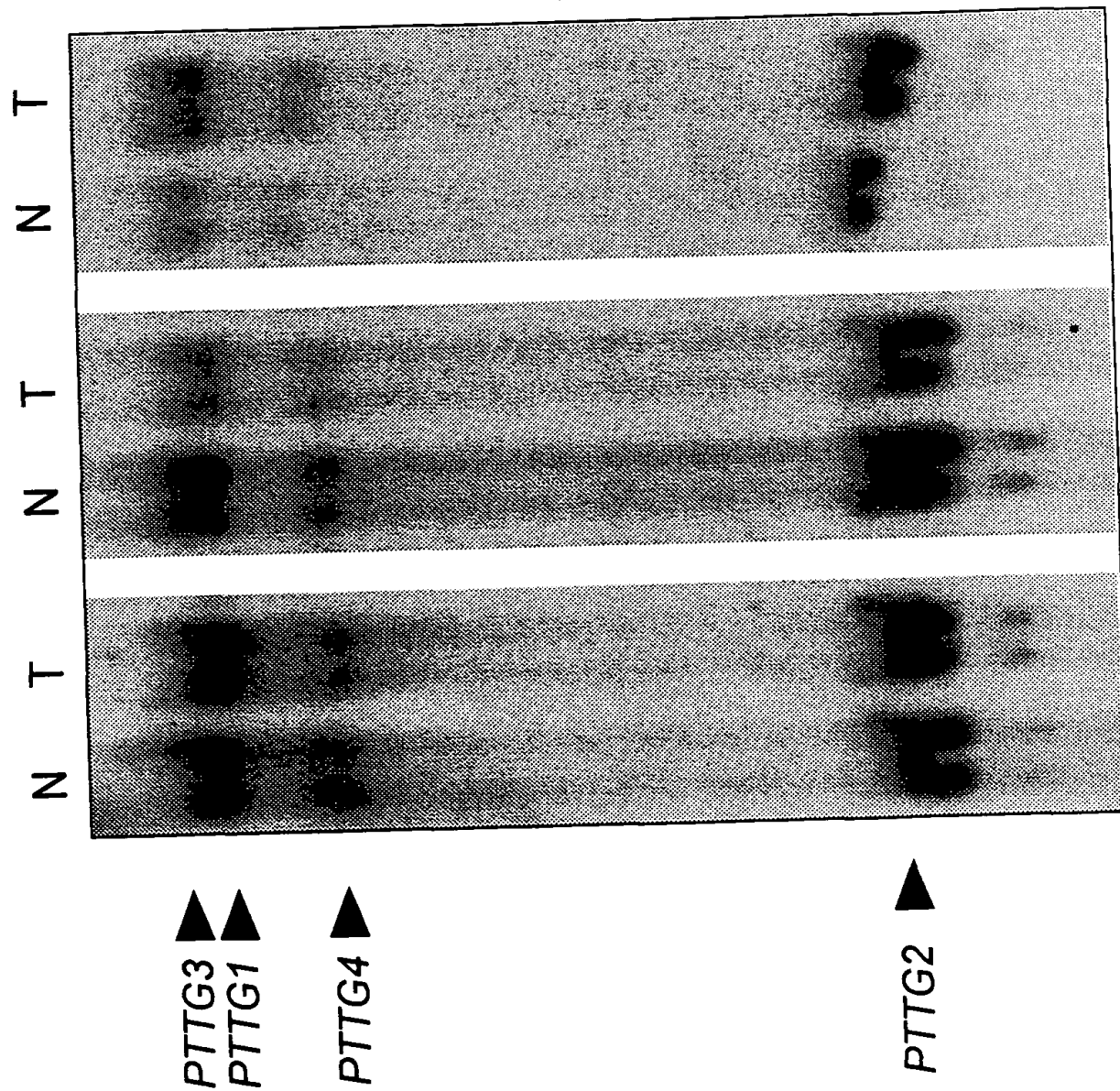


Figure 30B

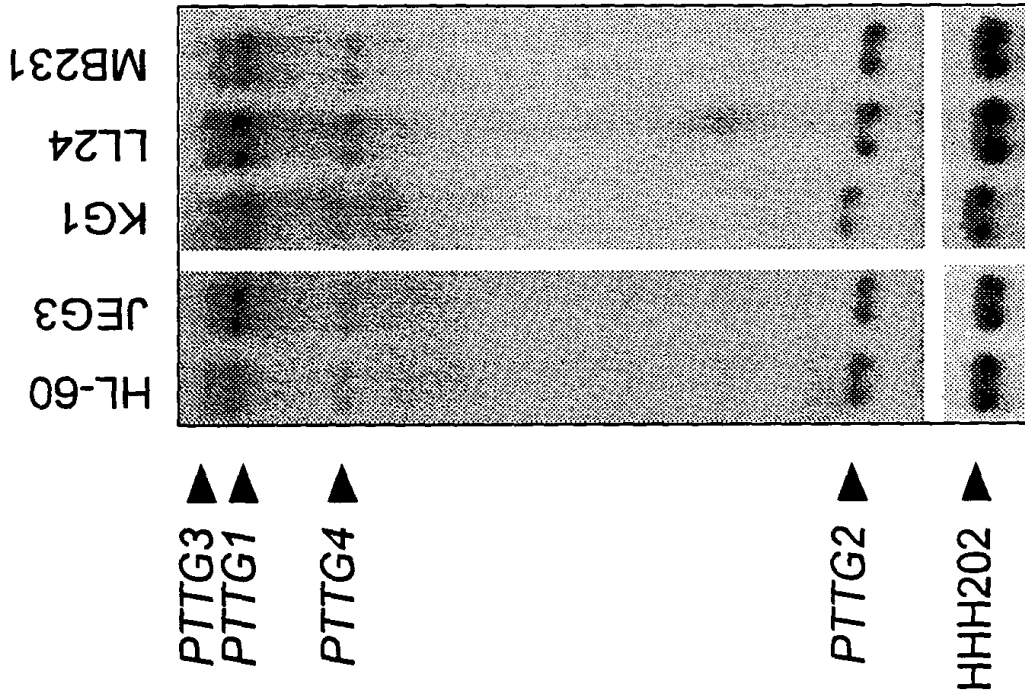


Figure 30C

09854325-051101

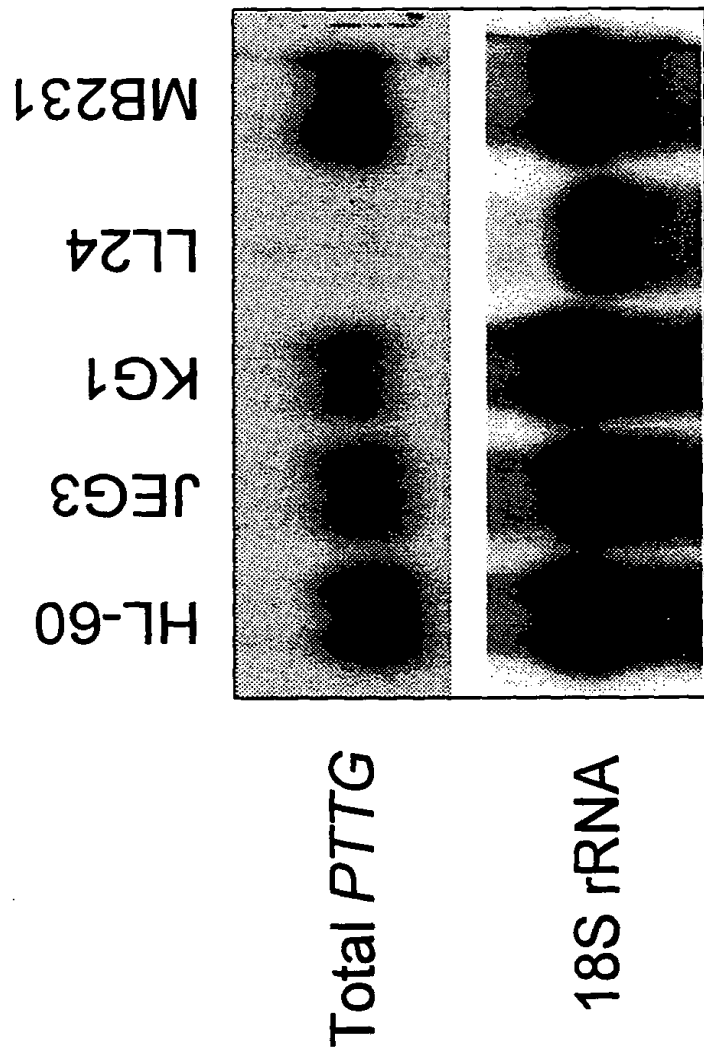
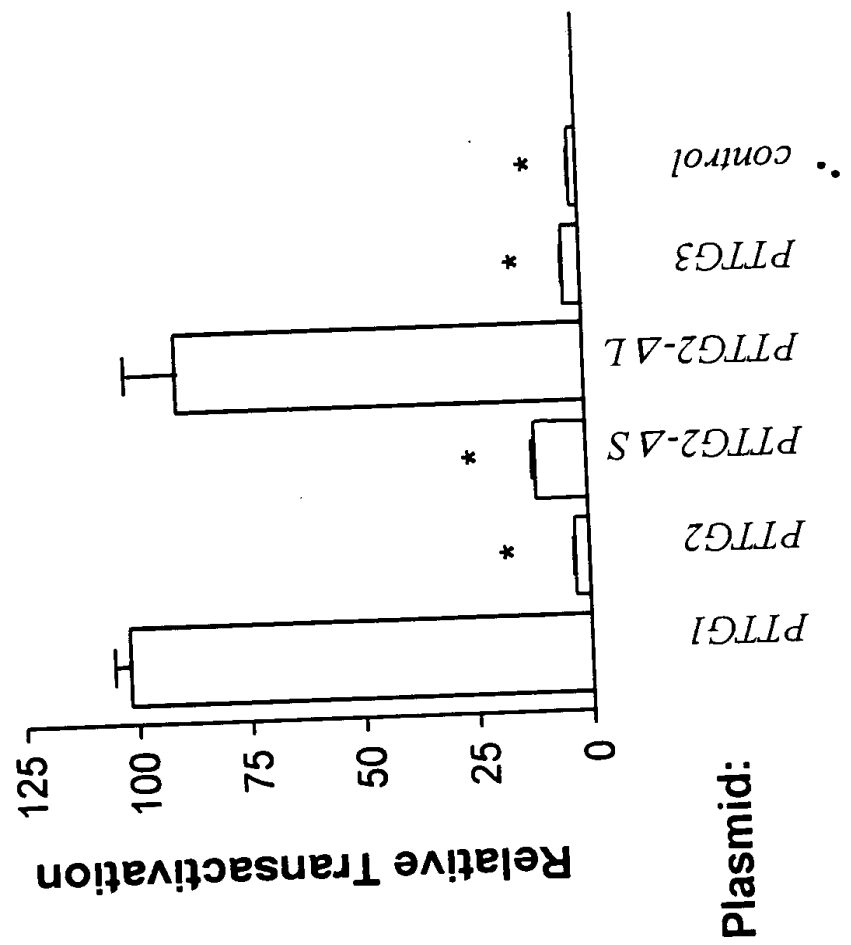


Figure 30D

Figure 31



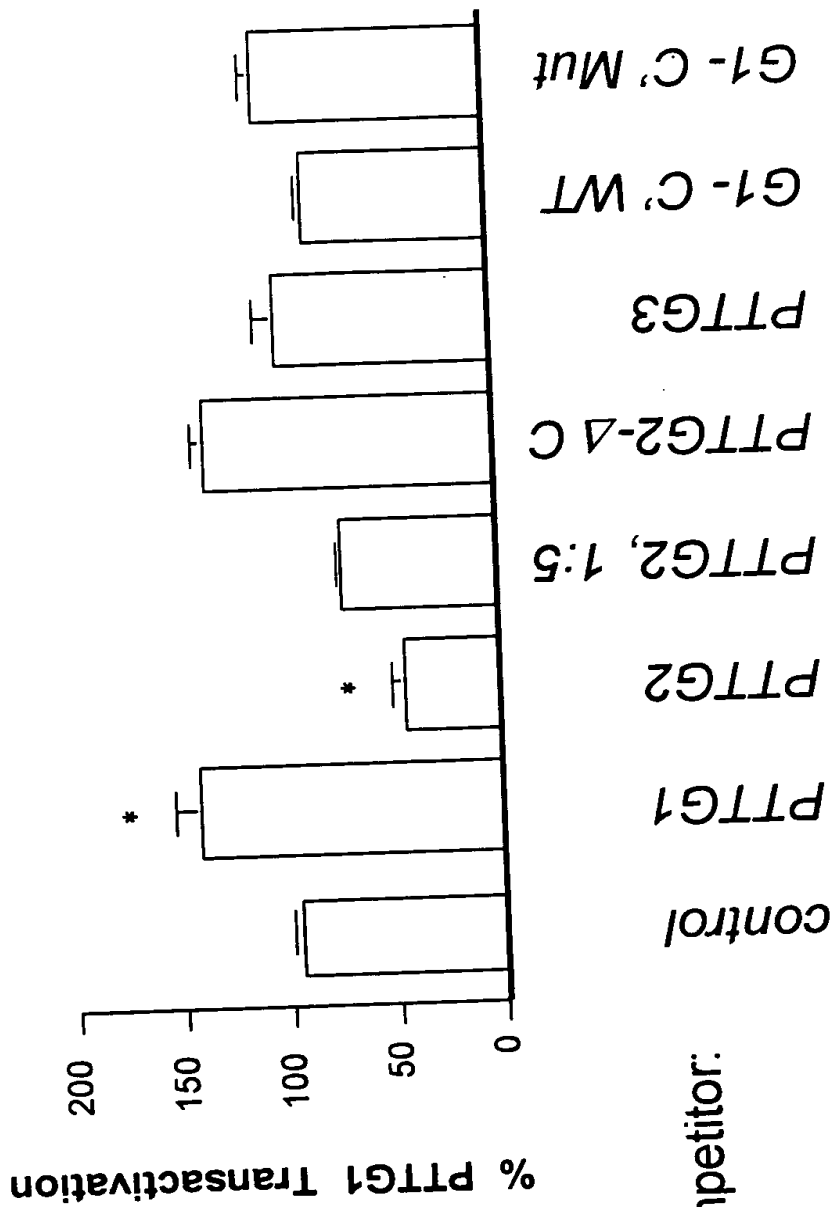


Figure 32A

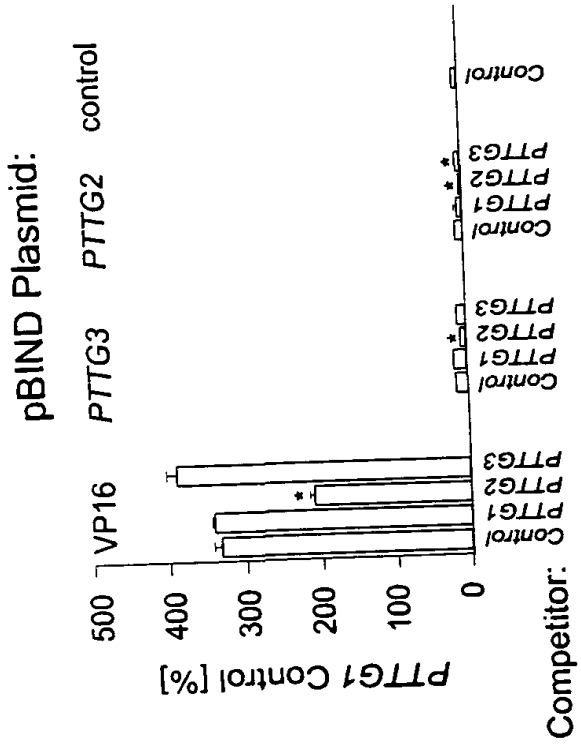


Figure 32B